Martin Fürer Institut für Theoretische Informatik, ETHZ Visiting from Pennsylvania State University

Joint work with David P. Jacobs an Vilmar Trevisan

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- -Introduction
 - └─ Tree-width and clique-width



- The subject started with Gaussian elimination for sparse matrices.
- Want to minimize fill-in.
- Intensive study of tree-width by Robertson and Seymour.
- Existence of polynomial time algorithms proved, without providing such algorithms.
- Minor closed classes of graphs. (A minor is obtained by vertex and edge deletions and edge contractions.)

- Introduction

└─ Tree-width and clique-width

Why tree-width?

- Many combinatorial graph problems are NP-hard.
- Usually, they are easy for trees.
- One wants to extend feasibility to a somewhat more general classes of graphs.
- The tree-width measures similarity to trees.
- Low tree-width usually implies efficient algorithms.

-Introduction

└─ Tree-width and clique-width

Tree-width

- Defined with the help of a tree-decomposition.
- Tree-width tw(G): Smallest k, having a tree decomposition with all bags of size ≤ k + 1.
- There are many efficient algorithms for graphs of small tree-width.
- Courcelles (1993) theorem: O(f(k) n) time algorithms (linear FPT) for all Monadic Second Order properties of vertices and edges.

- Introduction

└─ Tree-width and clique-width

Wanted: More powerful graph classes

- Bounded tree-width graphs are sparse.
- Most problems are easy for simple dense graphs like K_n or K_{pq}.
- Extend to a nice class?
- Intuitive property: Easily formed by adding all edges between two sets of vertices.
- Clique-width measures the complexity of such constructions.
- It is a more recently defined width parameter.

Locating the eigenvalues for graphs of small clique-width

- Introduction

└─ Tree-width and clique-width

k-expression defining a labeled graph

• Label set $= [k] = \{1, 2, \dots, k\}.$

Operations:

- i(v) create vertex v with label i.
- $\eta_{i,j}$ create all edges between the vertices labeled *i* and the vertices labeled *j* (for $i \neq j$).
- $\rho_{i \rightarrow j}$ change all labels *i* to *j*.
 - \oplus disjoint union of two graphs
- At the end, forget the labels.
- Clique-width cw(G) = smallest number of labels that can produce G.
- E.g., a clique of any size has clique-width 2.

-Introduction

Locating eigenvalues with Sylvester's law of inertia

Locating eigenvalues

- The number of eigenvalues grater than c is equal to the number of positive eigenvalues of B = A c I.
- Thus the number of eigenvalues in an interval can be determined by computing the number of positive and negative eigenvalues of two matrices.
- For regular P, the matrix $P^T B P$ is congruent to B.
- Sylvester: Congruent matrices have the same number of positive (negative) eigenvalues.
- Transform *B* into a congruent diagonal matrix.

└─ The result

The result

Theorem

For graphs of bounded clique-width with a given k-expression, the number of eigenvalues in an interval can be computed in linear time.

L The algorithm

└─ The shape of the matrix

The shape of the matrix B_q (congruent to B_0)



└─ The algorithm

- Methodology

Methodology

- For clique-width k, we focus on a small $p \times p$ submatrix (square box), where $p \le 2k$.
- Idea: We aim at the submatrix containing the rows and columns of one representative vertex per label.
- The algorithm uses the parse tree of a k-expression, and performs matrix operations corresponding to the operations in the k-expression.
- We only use congruence operations that add a multiple of a row and column to an other row and column respectively.

- └─ The algorithm
 - -The simple operations

The simple operations



- i(v): Focus on the 1×1 square box of vertex v.
- \blacksquare \oplus : Focus on square box around the two given square boxes.
- $\eta_{i,j}$: Insert some 1's in the square box.
- $\rho_{i \to i}$ record the new label.

└─ The algorithm

Congruency operations

Two vertices, say 1 and *i* have the same label: First step

$$\begin{pmatrix} \begin{bmatrix} a_0 & b_0 & \cdots & & \beta_1 \\ b_0 & c_0 & \cdots & & & \beta_2 \\ \vdots & \vdots & & & \vdots \\ & & & & & \beta_p \\ & & & & & & \beta_p \\ & & & & & & & \beta_p \\ & & & & & & & & \beta_p \\ & & & & & & & & & \beta_p \\ & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & \beta_p \\ & & & & & & & & & & & & \beta_p \\ \end{array} \right)$$

- In pictures, we omit the already diagonalized part for simplicity.
- The box represents the interesting submatrix.
- $\vec{0}$ and the β_i are (currently uninteresting) row-vectors.
- We show the first step when vertices 1 and *i* have the same label, implying $\beta_1 = \beta_i$.
- Subtract row/column i from row/column 1.

The algorithm

└─ Congruency operations





- Case 1: $a \neq 0$: Clear first row and column except for a in diagonal.
- Case 2: Top row is all 0: 0 in diagonal.
- Case 3: a = 0, top row not all 0, w.l.o.g. $b \neq 0$.

Locating the eigenvalues for graphs of small clique-width

- └─ The algorithm
 - Congruency operations

The interesting Case 3

$$\begin{pmatrix} \begin{matrix} 0 & b & \cdots & & \vec{0} \\ b & c & \cdots & & & \beta_2 \\ \vdots & \vdots & & & & \vdots \\ & & & & & \beta_p \\ \vec{0}^T \beta_2^T \dots \beta_p^T \end{pmatrix} \Rightarrow \begin{pmatrix} \begin{matrix} 0 & b & \vec{0} & & \vec{0} \\ b & c & \cdots & & & \beta_2 \\ & & & & & \vec{0}^T & \vdots & & & \vdots \\ & & & & & & & \beta_p' \\ \vec{0}^T \beta_2^T \dots \beta_p'^T & & \end{pmatrix}$$

• Only Case 3 (a = 0) needs more work.

- Add appropriate multiples of row/column 2 to clear column/row 1 after position 2.
- For some constants c_3, \ldots, c_p , $(\beta_3, \ldots, \beta_p)$ gets replaced by $(\beta'_3, \ldots, \beta'_p) = (\beta_3 + c_3\beta_2, \ldots, \beta_p + c_p\beta_2).$

Locating the eigenvalues for graphs of small clique-width

- └─ The algorithm
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The interesting Case 3



- Add multiples of row/column 1 to clear column/row 2 after position 2.
- Add (1 c/b)/2 times row/column 1 to row/column 2. Now c is b.
- Add multiples of row/column 1 to all rows/colums after 2 to clear column/row 2 after position 2.
- Subtract row/column 2 from row/column 1.
- Now the first two rows and columns are diagonalized.

The algorithm

Congruency operations

Complication

$$B_{q} = \begin{pmatrix} \begin{bmatrix} -b & 0 & \vec{0} \\ 0 & b & \vec{0} \\ \vec{0}^{T} & \vec{0}^{T} \\ & & \\ & & \\ \hline \vec{0}^{T} & \vec{0}^{T} \\ & & \\ & & \\ \hline \vec{0}^{T} & \vec{0}^{T} \\ & & \\ \end{pmatrix}$$

- β_2 has disappeared as its own row vector. It is unrepresented by the matrix B_q
- It shows up in other rows as $\beta'_i = \beta_i + c_i\beta_2$ for i > 2.
- Thus, as more entries of β_2 get known, β'_i has to be updated.
- When another vertex later shows up with label 2, and thus with row vector β₂, then β_i can be reconstructed by subtracting c_iβ₂ from β'_i.

└─ The algorithm

Congruency operations

More complications

- At the beginning, we assumed two vertices, 1 and *i* have the same label and hence $\beta_1 = \beta_i$. Thus $\vec{0}$ has been created by subtraction.
- In reality, we subtact β'_i from β'_i .
- Instead of 0, we obtain a linear combination of the vectors of the unrepresented vertices.
- Instead of aiming at having just one row (and column) per label, we represent up to u additional rows, where u is the number of unrepresented vertices (which is less than the clique-width k).
- The additional rows are intuitively close to 0. They are just linear combinations of the vectors of the unrepresented vertices.
- If one more additional row is obtained, then 0 can be linearly combined, and we can continue as in the typical case where the earlier Case 3 never happened.

└─ The algorithm

Congruency operations

The end

Thank you!